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172	280.5	305.5	357	380	383	388	392	395	395	397	397	400	400
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ALIGNMENTS

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ID = CD4 H
AC P0173
DT 25-JA
DT 01-NO
DT 01-NO
DT 7-Cel
DE T4-LCe
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P01730;
MEDLINE-22388257; PubMed=12477932; DOI=10.1073/pnas.242603899; Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G., Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D. Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K., Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F., Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L., Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T. Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A., AND VARIANT TRP-265.

MEDLINE=91216786; PubMed=1708753; DOI=10.1016/0198-8859(91)90077-M;

MODELINE=91216786; PubMed=1708753; DOI=10.1016/0198-8859(91)90077-M;

Hodge T.W., Sasso D.R., McDougal J.S.;

"Humans with OKT4-epitope deficiency have a single nucleotide base change in the CD4 gene, resulting in substitution of TRP240 for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=85254948; PubMed=2990730;
Maddon P.J., Littman D.R., Godfrey M., Maddon D.E., Chess L., Axel R.
"The isolation and nucleotide sequence of a cDNA encoding the T cell
surface protein T4: a new member of the immunoglobulin gene family.";
Cell 42:93-104(1985).
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Ansari-Lari M.A., Muzny D.M., Lu J., Lu F., Lilley C.E., Spanos S.
Malley T., Gibbs R.A.;
"A gene-rich cluster between the CD4 and triosephosphate isomerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Littman D.R., Maddon P.J., "Corrected CD4 sequence."; Cell 55:541-541(1988).
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Mammalia; Eutheria; Primates;
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                     change in the CD4 gene, ARG240.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             genes at human chromosome 12p13.";
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Littman D.R., Maddon P.J., Axel R.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Name=CD4;
Homo sapiens (Human)
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                                                                                                                                                                                                                                                                                          TISSUE=Pancreas
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(Rel. 09, Last sequence update)
(Rel. 46, Last annotation update)
ace glycoprotein CD4 precursor (T-cell surface antigen
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Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J., Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H., Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W., Villalon D. K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A., Villalon D. K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A., Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A., Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G., Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C., Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C., Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Shalter A., Schein J.E., Jones S.J.M., Marra M.A.; Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.; "Generation and initial analysis of more than 15,000 full-length huma and mouse cDNA sequences."; c. 00.1600-16000-160001
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MEDLINE-90078232; PubMed=2592374;

Carr S.A., Hemling M.E., Folena-Wasserman G.,

Barr J.R., Huddleston M.J., Taylor P.;

"Protein and carbohydrate structural analysis

CD4 receptor by mass spectrometry.";

J. Biol. Chem. 264:21286-21295(1989).
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"Cloning and sequences of primate CD4 molecules: diversity of the cellular receptor for simian immunodeficiency virus/human immunodeficiency virus.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PubMed=15340161; DOI=10.1110/ps.04682504; Zhang Z., Henzel W.J.; "Signal peptide prediction based on analy verified cleavage sites.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "Nef induces CD4 endocytosis: requirement for a critical ("notif in the membrane-proximal CD4 cytoplasmic domain."; Cell 76:853-864(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=93049640; PubMed=1425921;
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SER-433; 438-LEU-LEU-439 AND
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MEDLINE=91061881; PubMed=1701030; DOI=10.1038/348411a0;

Mang J., Yan Y., Garrett T.P., Liu J., Rodgers D.W., Garlick Tarr G.E., Husain Y., Reinherz E.L., Harrison S.C.;

"Atomic structure of a fragment of human CD4 containing two immunoglobulin-like domains.";

Nature 348:411-418(1990).
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MEDLINE-91061882; PubMed=2247146; DOI=10.1038/348419a0;
Ryu S.-E., Kwong P.D., Truneh A., Porter T.G., Arthos J.,
Rosenberg M., Dai X., Xuong N.-H., Axel R., Sweet R.W.,
PALMITOYLATION.
MEDLINE=92317088;
Crise B., Rose J.1
                                                                                                  MEDLINE=97311402; PubMed=9168119;
Wu H., Kwong P.D., Hendrickson W.
"Dimeric association and segmenta
                                                                                                                                                                          Nature 348:419-426(1990)
[12]
                                                                                                                                                                                                                                            Rosenberg M., Dai
Hendrickson W.A.;
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                                                                                                                                                        X-RAY CRYSTALLOGRAPHY (3.9
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                                                                       387:527-530(1997)
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     Rose J.K.;
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                         PubMed=1618861;
                                                                                                                            Hendrickson W.A.;
                                                                                                          segmental
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PIR; A90072; RWHUT4.

PIR; A90072; RWHUT4.

PDB; 1CDH; X-ray; @=26-203.

PDB; 1CDL; X-ray; @=26-203.

PDB; 1CDU; X-ray; @=26-203.

PDB; 1CDU; X-ray; @=26-203.

PDB; 1CDU; X-ray; @=26-210.

PDB; 1G9N; X-ray; C=26-210.

PDB; 1G9N; X-ray; C=26-210.

PDB; 1G14; X-ray; C=26-210.

PDB; 1G2; X-ray; C=26-203.

PDB; 1MIA; X-ray; D=26-203.

PDB; 1MIA; NMR; A=421-445.

PDB; 1MIA; X-ray; A/B=26-388.

PDB; 1MIC; X-ray; A/B=26-388.
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EMBL;
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Pfam; PF00047; 1g; 2.
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InterPro; IPR000973;
InterPro; IPR007110;
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TRANSMEM
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                                                                                                                                                                                                                                                        GO:0005886; C:plasma membrane; TAS.
GO:0042101; C:T-cell receptor complex; NAS.
GO:0042102; F:coreceptor activity; NAS.
GO:0015026; F:coreceptor activity; NAS.
GO:0015029; F:nHc class II protein binding; NAS.
GO:0042289; F:transmembrane receptor activity; TAS.
GO:004955; P:transmembrane receptor activity; TAS.
GO:0006955; P:mmune response; NAS.
GO:0045086; P:positive regulation of interleukin-2 biosyn.
GO:0045086; P:T-cell differentiation; NAS.
GO:0045058; P:T-cell selection; NAS.
GO:0045059; P:transmembrane receptor protein tyrosine kin.
GO:007169; P:transmembrane receptor protein tyrosine kin.
                                                                                                                                                                                                                                                                                                                                                                                                                                         186940;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                M12807; AAA35572.1; -. U47924; AAB51309.1; -. M35160; AAA16069.1; -. BC025782; AAH25782.1;
                                                                                                                            ; PR00692; CD4TCANTI(
SM00406; IGv; 1
B; PS50835; IG_LIKE;
                  26
26
397
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cy virus receptor.";
267:13593-13597(1992)
                                                                    Repeat; Signal;
                                                                                                                                                              CD4TCANTIGEN
     458
418
418
                                                                                                                                                                                                                CD4 TCAg.
Ig-like.
     T-cell surface (Extracellular (Epotential. Cytoplasmic (Pot
                                                                                       T-cell;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (See http://www.isb-sib.ch/announce/
                                                                                         Transmembrane
            (Potential)
                                           glycoprotein (Potential).
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                                                                                                            Palmitate;
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RESULT
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Best Local Sim
Matches 199;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PANTR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-APR-1990
01-APR-1990
05-JUL-2004
                                                                                                      This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Event the Event of Event of the Event of Event of the Event of Event of Event of the Event of 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fomsgaard A., Hirsch V.M., Johnson P.R.; "Cloning and sequences of primate CD4 molecules: cellular receptor for simian immunodeficiency vi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CD4 PANTR
P16004;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DOMAIN
              EMBL; M31135; AAA35407.1; -.
EMBL; X73323; CAA51749.1; -.
PIR; B32722; RWCZT4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "A CD4 domain important for HIV-mediated outside the virus binding site.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=9598;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pan troglodytes (Chimpanzee).
Eukaryota; Metazoa; Chordata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      T-cell surface T4/Leu-3).
                                                                                                                                                                                                                                                                                                                                                                                                                                 immunodeficiency virus.";
Eur. J. Immunol. 22:2973-2981(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=93049640; PubMed=1425921;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE OF 26-424 FROM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Camerini
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE=90182664; PubMed=2107024; DOI=10.1016/0092-8674(90)90089-W;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Name=CD4
                                                                                                                                                                                                                                                                                    FUNCTION: Accessory protein for MHC class-II antigen/T-cell receptor interaction. May regulate T-cell activation. SUBUNIT: Associates with p56-lck (By similarity). SUBCELLULAR LOCATION: Type I membrane protein. SIMILARITY: Contains 3 immunoglobulin-like C2-type domains. SIMILARITY: Contains 1 immunoglobulin-like V-type domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      121
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           D., Seed B.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MNRGVPFRHLLLVLQLALLPAATQGNKVVLGKKGDTVELTCTASQKKSIQFHWKNSNQIK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TWTCTVLQNQKKVEFKIDIV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ILGNOGSFLTKGPSKLNDRADSRRSLWDQGNFPLIIKNLKIEDSDTYICEVEDQKEEVQL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (Rel. 14, Created)
(Rel. 14, Last sequence update)
(Rel. 44, Last annotation update)
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99.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 1023; Di
Pred. No. 3.5e
0; Mismatches
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Ig-like
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Pan.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C2-type 1.
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3.5e-77;
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                                                                                                                                . Usage by and fc http://www.isb-sib.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                es: diversity virus/human
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Best Local &
Matches 195
                                                                   MACFA STANDARD; PRT; 458 AA.
PY9185;
15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
T-cell surface glycoprotein CD4 precursor (T-T4/Leu-3).
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CONFLICT
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InterPro;
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DOMAIN
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Macaca fascicularis
Eukaryota; Metazoa;
Mammalia; Eutheria;
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                                                       Name=CD4;
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Palmitate; Repeat; Signal;
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GO:0015026; F:coreceptor activity; ISS.
[GO:0042209; F:Mfc class II protein binding; ISS.
GO:0006955; P:immune response; ISS.
GO:0045086; P:positive regulation of interleukin-2
GO:0045086; P:T-cell differentiation; ISS.
GO:0045088; P:T-cell selection; ISS.
GO:0045088; P:T-cell selection; ISS.
GO:007159; P:transmembrane receptor protein tyrosi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; IPR008424; CD2.
; IPR000973; CD4 TCAg.
; IPR007110; Ig-Tike.
; IPR003596; Ig_v.
                                                                                                                                                                                                                                                                                                                TWTCTVLQNQKKVEFKIDIV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MNRGVPFRHLLLVLQLALLPAATQGKKVVLGKKGDTVELTCTASQKKSIQFHWKNSNQTK
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26 125
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1191 KW;
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(Crab eating macaque) (Cynomolgus monkey).
Chordata, Craniata; Vertebrata; Euteleostomi;
Primates; Catarrhini; Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 97.1%;
97.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 response; Immunoglobulin domain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ig-like V-type.
Ig-like C2-type 1.
Ig-like C2-type 3.
Ig-like C3-type 3.
N-linked (GlcNAc...)
N-linked (GlcNAc...)
By similarity.
By similarity.
By similarity.
S-palmitcyl cysteine (S-palmitcyl cysteine (T-> N (in Ref. 2).
L-> S (in Ref. 2).
K-> N (in Ref. 2).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 999; DB
Pred. No. 3.6e
0; Mismatches
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Cytoplasmic
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Extracellular (Potential).
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T-cell surface
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similarity)
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Best Local (
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HSSP; P01730; IMBR.
GO; GO:00042101; C:T-cell recep
GO; GO:0015026; F:coreceptor a
GO; GO:0042289; F:MHC class II
GO; GO:0006955; P:immune respo
GO; GO:0045086; P:positive respo
GO; GO:0030217; P:T-cell diffe
GO; GO:0045058; P:T-cell selec
GO; GO:0007169; P:T-cell selec
GO; GO:0007169; P:transmembran
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NCBI_TaxID=9541;
[1]
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Submitted (FEB-1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              the European Bioinformatics Institute. The use by non-profit institutions as long modified and this statement is not removed.
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PROSITE; PS50835; IG_LIKE; 1.
Glycoprotein; Immune_response;
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InterPro; IPR00973; CD4 TCAg.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
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Palmitate; Re
                                                                                                                                                                                                                                                                                                                                                                                                           SIGNAL
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                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sumi M., Yabe M., Yamada Y.K.;
mitted (FEB-1997) to the EMBL/GenBank/DDBJ databases.
mUNCTION: Accessory protein for MHC class-II antigen/T-cell
receptor interaction. May regulate T-cell activation.
SUBUNIT: Associates with p56-lck (By similarity).
SUBURLARIFY: Contains 3 immunoglobulin-like C2-type domains.
SUBULARITY: Contains 1 immunoglobulin-like V-type domains.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GO:0042101; C:T-cell receptor complex; ISS.
GO:0042102; F:coreceptor activity; ISS.
GO:0015026; F:coreceptor activity; ISS.
GO:0042289; F:MHC class II protein binding; ISS.
GO:0006955; P:immune response; ISS.
GO:0005056; P:positive regulation of interleukin-2 biosyn.
GO:00030217; P:T-cell differentiation; ISS.
GO:0045058; P:T-cell selection; ISS.
GO:0007169; P:transmembrane receptor protein tyrosine kin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SWISS-PROT entry is copyright. It is produced through a collaboration een the Swiss Institute of Bioinformatics and the EMBL outstation - European Bioinformatics Institute. There are no restrictions on its by non-profit institutions as long as its content is in no way
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; PR00692; CD4;
SM00406; IGV;
61
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                                                                                                                          Similarity
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                                                      ILGNQGSFLTKGPSKLNDRADSRRSLWDQGNFPLIIKNLKIEDSDTYICEVEDQKEEVQL
 ILGIQGSFLTKGPSKLSDRADSRKSLWDQGCFSMIIKNLKIEDSDTYICEVENKKEEVEL
                                                                                                                                                                                                                                                                                                                                                                                                                        Repeat;
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                                                                                                                                                                                                                                                                                             Signal;
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                                                                                                                                                                Potential.

Cytoplasmic (Potential).

Ig-like V-type.

Ig-like C2-type 1.

Ig-like C2-type 2.

Ig-like C2-type 2.

Ig-like C2-type 2.

In-linked (GlcNAc. . .) (Potential).

N-linked (GlcNAc. . .) (By similarity).

N-linked (GlcNAc. . .) (By similarity).

Sysimilarity.

By similarity.

By similarity.

By similarity.

By similarity.

S-palmitoyl cysteine (By similarity).

S-palmitoyl cysteine (By similarity).

S-palmitoyl cysteine (By similarity).
                                                                                                              12;
                                                                                                              Score 912; DB
Pred. No. 6.8e
L2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                            By similarity.
T-cell surface glycoprotein CD4.
Extracellular (Potential).
                                                                                                                                                                                                                                                                                                                                                                                                                          -cell; Transmembrane.
                                                                                                                                                                                                                                                                                                                                                                                                                                     Immunoglobulin domain; Lipoprotein;
                                                                                                                           DB 1;
.8e-68;
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(By similarity).
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CD4 MACFU
P79184;
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Hashinoto O., Tatsumi M.;
Submitted (FEB-1997) to the EMBL/GenBank/DDBJ databases.
Submitted (FEB-1997) to the EMBL/GenBank/DDBJ databases.
-I- FUNCTION: Accessory protein for MHC class-II antigen/T-cell receptor interaction. May regulate T-cell activation.
-I- SUBUNIT: Associates with p56-lck (By similarity).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Macaca fuscata fuscata (Japanese macaque).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               P79188;
15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
05-JUL-2004 (Rel. 44, Last annotation update)
                                                                                                                                                                                                                                                                                                     EMBL; D63348; BAA09672.1; -.

HSSP; P01730; IMBR.
G0; G0:0042101; C:T-cell receptor complex; ISS.
G0; G0:0042204; F:Coreceptor activity; ISS.
G0; G0:0015026; F:coreceptor activity; ISS.
G0; G0:0042289; F:MHC class II protein binding; ISS.
G0; G0:0042289; F:immune response; ISS.
G0; G0:0045086; P:positive regulation of interleukin-2 biosyn.
G0; G0:0045086; P:T-cell differentiation; ISS.
G0; G0:0045089; P:T-cell differentiation; ISS.
G0; G0:0045089; P:T-cell selection; ISS.
G0; G0:0007159; P:T-cell selection; ISS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cercopithecinae; Macaca.
NCBI_TaxID=9543;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial
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 DOMAIN
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                            TRANSMEM
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                                                                                                                                                                                                Pfam; PF00047; ig; 2. PRINTS; PR00692; CD4T SMART; SM00406; IGV;
                                                                                                                                                                                                                                         InterPro; IPR008424; CD2.
InterPro; IPR000973; CD4 TCAg.
InterPro; IPR007110; Ig-Tike.
InterPro; IPR003596; Ig_v.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          entities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                                                   Glycoprotein; Immune Palmitate; Repeat; Si
                                                                                                                                                                                SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
                                                                                                         DOMAIN
                                                                                                                                        SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SUBCELLULAR LOCATION: Type I membrane protein.
SIMILARITY: Contains 3 immunoglobulin-like C2-type domains.
SIMILARITY: Contains 1 immunoglobulin-like V-type domain.
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317
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                                                                                                                                                   Signal;
                                                                                                                                                                    response; Immunoglobulin domain; Lipoprotein;
 T-cell surface glycoprotein C
Extracellular (Potential).
Potential.
Cycoplasmic (Potential).
Ig-like V-type.
Ig-like C2-type 1.
Ig-like C2-type 2.
Ig-like C2-type 3.
N-linked (GlcNAc. . .) (Poten
                                                                                                                                                         T-cell;
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                                                                                                                           By similarity. T-cell surface
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                                                                                                                                                         Transmembrane.
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ID CD4 M
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DT 01-Ap
DT 28-FE
DT 14/Le
GN Macac
OC Camer
OC CAMEN
OC CAMEN
RY MEDLI
RN 13
RP SEQUE
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Best Local S
Matches 175
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CD4 MACMU
CD5 MACMU
P16003; Q29617;
01-APR-1990 (Rel. 14, Created)
28-FBB-2003 (Rel. 41, Last sequence update)
05-UTL-2004 (Rel. 44, Last annotation update)
05-UTL-2004 (Rel. 44, Created)
05-UTL-2004 (Rel. 44, Created)
05-UTL-2004 (Rel. 47, Last annotation update)
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SEQUENCE OF 107-192 FROM N.A. MEDLINE=98320644; PubMed=9656488; Harris E.E., Disotell T.R.; "Nuclear gene trees and the phylogenetic mangabeys (Primates: Papionini).";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A. MEDLINE=90182664; PubMed=2107024;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Macaca mulatta (Rhesus macaque).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Primates; Catarrhini; Cercopithecidae;
                                                                                                                           Fomsgaard A., Hirsch V.M., Johnson P.R.; "Cloning and sequences of primate CD4 molecular cellular receptor for simian immunodeficiency immunodeficiency virus."; "Lur. J. Immunol. 22:2973-2981 (1992).
                                                                                                                                                                                                                                                                                                                         TISSUE=Thymocytes;
Hashimoto O., Tatsumi M.;
"Molecular cloning and exp
Submitted (JUN-1995) to the
                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Camerini D., Seed B.;
"A CD4 domain important for HIV-mediated syncytium formation lies outside the virus binding site.";
Cell 60:747-754(1990).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cercopithecinae; Macaca.
NCBI_TaxID=9544;
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                                                                                                                                                                                                                                      MEDLINE=93049640; PubMed=1425921;
                                                                                                                                                                                                                                                             TISSUE=Blood;
                                                                                                                                                                                                                                                                                 SEQUENCE OF 28-424 FROM N.A.
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S-palmitoyl cysteine (()
76B3E7EP08185535 CRC
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No. 1e-67;
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                                                                                                                                                                          molecules: diversity of the 
ficiency virus/human
                        relationships
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PRINTS; PR00652; CD4TCANTIGEN.
SMART; SM00406; IGV; 1.
PROSITE; PS50835; IG_LIKE; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GG; GO:0007169; P:transmembrane InterPro; IPR008424; CD2. InterPro; IPR00873; CD4 TCA9. InterPro; IPR007110; Ig-Tike. InterPro; IPR003596; Ig_v.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HSSP;
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EMBL; D63347;
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                                                                                              SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                  DOMAIN
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                                                                                                                                                                                                                                                                                                                                                                                                      Glycoprotein; Immu
Palmitate; Repeat;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GO:0042101; C:T-cell receptor complex; ISS.
GO:0015026; F:coreceptor activity; ISS.
GO:0042289; F:MHC class II protein binding; ISS.
GO:0006955; P:immune response; ISS.
GO:0045086; P:positive regulation of interleukin-2
GO:0030217; P:T-cell differentiation; ISS.
GO:0045088; P:T-cell selection; ISS.
GO:0045088; P:T-cell selection; ISS.
GO:0045088; P:T-cell selection; ISS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FUNCTION: Accessory protein for MHC class-II antigen/T-cell receptor interaction. May regulate T-cell activation. SUBUNIT: Associates with p56-lck (By similarity).

SUBCELLULAR LOCATION: Type I membrane protein.

SIMILARITY: Contains 3 immunoglobulin-like C2-type domains.

SIMILARITY: Contains 1 immunoglobulin-like V-type domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Biol. Evol. 15:892-900(1998).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AF057385; AAC25129.1; -. P01730; 1WBR.
                                                            Similarity
             MNRGVPFRHLLLVLQLALLPAATQGNK/VVLGKKGDTVBLTCTASQKKSIQFHWKNSNQIK
MRGIPFRHLLVLQLALLPAVTQGKKVVLGKKGDTVELTCNASQKKNTQFHWKNSNQIK
                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                      Signal; T-cell; Transmembrane
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418
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422
422
169
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                                                            87.9%;
87.0%;
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By similarity.
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S-palmitoyl cysteine (
S-palmitoyl cysteine (
N-> T (in Ref. 1).
L-> S (in Ref. 2).
L-> L (in Ref. 2).
K-> N (in Ref. 2).
K-> N (in Ref. 2).
S-> P (in Ref. 2).
S-> P (in Ref. 3).
S-> P (in Ref. 3).
A-> T (in Ref. 3).
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Ig-like C2-type 1.
Ig-like C2-type 2.
Ig-like C2-type 3.
N-linked (GlcNAc...
N-linked (GlcNAc...
                                               Score 904; DE
Pred. No. 3.2e
L3; Mismatches
                                                                                                                                                                                                                                                                                                                                          Potential.
Cytoplasmic (Potential).
                                                                                                                                                                                                                                                                                                                                                                  T-cell surface glycoprotein 
Extracellular (Potential).
                                                                                                                                                                                                                                                        similarity.
                                                           DB 1;
.2e-67;
                                                                      Length 458;
                                                                                                 CRC64;
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Q08340; P79196;
01-FEB-1995 (Rel. 31, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
T-cell surface glycoprotein CD4 precursor (T-cell
T4/Leu-3).
InterPro; IPR008424; CD2.
InterPro; IPR008973; CD4 TCA9.
InterPro; IPR007110; Ig-Tike.
InterPro; IPR003596; Ig_v.
Pfam; PF00047; 19; 2.
PRINTS; PR00692; CD4TCANTIGEN.
                                                                                                                                                                                                                                                                                                                                          Pomsgaard A., Hirsch V.M., Johnson P.R.;

Poloning and sequences of primate CD4 molecules: diversity of the collular receptor for simian immunodeficiency virus/human immunodeficiency virus/human immunodeficiency virus.";

Eur. J. Immunol. 2::2973-2981(1992).

Eur. J. Immunol. 2::2973-2981(1992).

Function: Accessory protein for MRC class-II antigen/T-cell receptor interaction. May regulate T-cell activation.

Function: Function: Type I membrane protein.

SUBCELULAR LOCATION: Type I membrane protein.

SIMILARITY: Contains 1 immunoglobulin-like V-type domains.

SIMILARITY: Contains 1 immunoglobulin-like V-type domains.
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Mammalia; Eutheria;
Cercopithecinae; Mac
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Macaca nemestrina (Pig-tailed
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Name=CD4;
                                                                                                                                                                                                                            This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE OF 28-424 FROM N.A.
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Hashimoto O., Tatsumi
Submitted (FEB-1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=9545;
                                                                                                                                                                                                     EMBL;
                                                                              GO:0042101, C:T-cell receptor complex; ISS.
GO:0015026; F:coreceptor activity; ISS.
GO:0042289; F:MHC class II protein binding; ISS.
GO:0006955; P:immune response; ISS.
GO:0045086; P:positive regulation of interleukin-2
GO:0045086; P:T-cell differentiation; ISS.
GO:0045088; P:T-cell selection; ISS.
GO:0045088; P:T-cell selection; ISS.
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Craniata; Vertebrata; Euteleostomi;
Catarrhini; Cercopithecidae;
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pROSITE; pS50835; IG_LIKE; 1.
glycoprotein; Immune response
palmitate; Repeat; Signal; T-
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Q08338; O02805; O77593; Q28217;
Q08338; O02805; O77593; Q28217;
Q1-FEB-1995 (Rel. 31, Created)
29-MAR-2004 (Rel. 43, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
T-cell surface glycoprotein CD4 precursor (T-cell T4/Leu-3).
                                                                                                                                                                                                                            Cercopithecus aethiops (Green m
Eukaryota; Metazoa; Chordata; C
Mammalia; Butheria; Primates; C
Cercopithecinae; Cercopithecus.
NCBI TaxID=9534;
                                                                                        "Molecular
Submitted (
                                                                                                                                                                                                                                                                                                                                                                                             Name=CD4;
SEQUENCE OF 28-424
TISSUE=Blood;
                                                                                        Hashimoto O., Tatsumi M. "Molecular cloning and e Submitted (JUL-1996) to
                                                                                                                                                                                SEQUENCE
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ignal; T-cell; Transmembrane
                                         FROM N.A.
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87.0%;
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Cytoplasmic (Potential Ig-like V-type. 1
Ig-like C2-type 1.
Ig-like C2-type 2.
Ig-like C2-type 3.
Ig-like C3-type 3.
N-linked (GlcNAc. ..)
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By similarity.
By similarity.
By similarity.
S-palmitcyl cysteine (S-palmitcyl cysteine (D-palmitcyl Cysteine (Cysteine 
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Pred. No. 3.8e-67;
4; Mismatches 12
                                                                                                    ssion of african green monke EMBL/GenBank/DDBJ databases
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                                                                                                                                                                                                                                                                                                           Catarrhini; Cercopithecidae
                                                                                                                                                                                                                                                                                                                                                 monkey) (Grivet).
Craniata; Vertebrata;
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Fomsgaard A., Hirsch V.M., Johnson P.R.; "Cloning and sequences of primate CD4 molecules: diversity of the cellular receptor for simian immunodeficiency virus/human

MBDLINE=93049640; PubMed=1425921;

lmmunodeficiency virus.";
3ur. J. Immunol. 22:2973-2981(1992).

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GO; GO:0042101; C:T-cell receptor complex; ISS.

GO; GO:0015026; F:coreceptor activity; ISS.

GO; GO:0015026; F:coreceptor activity; ISS.

GO; GO:0042289; F:MHC class II protein binding; ISS.

GO; GO:0006955; F:immune response; ISS.

GO; GO:0006955; P:positive regulation of interleukin-2 biosyn.

GO; GO:0003217; F:T-cell differentiation; ISS.

GO; GO:0007169; P:T-cell selection; ISS.

GO; GO:0007169; P:transmembrane receptor protein tyrosine kin.
                                                                                                                                                                                                                                                                                                                                                                                                                                         EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the European Bioinformatics Institute. The use by non-profit institutions as long a modified and this statement is not removed. entities requires a license agreement (See lor send an email to license@isb-sib.ch).
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Harris E.B., Disotell T.R.;
"Nuclear reso
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Glycoprotein; Immune respons
Palmitate; Repeat; Signal; T
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InterPro; IPR000973; CD4 TCAg.
InterPro; IPR007110; Ig-Tike.
                                                                                                                                                                                                                                                                                                                                                                                                                            EMBL;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TISSUE-Peripheral blood; MEDLINE-98017879; PubMed-9379478;
                                                                                                                                                                            SIGNAL
                                                                                                                                                                                                                                  PRINTS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "Relation between phylogeny of African green monkey CD4 their respective simian immunodeficiency virus genes."; J. Med. Primatol. 26:120-128(1997).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE OF 28-424 FROM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FUNCTION: Accessory protein for MHC class-II antigen/T-cell receptor interaction. May regulate T-cell activation. SUBUNIT: Associates with p56-1ck (By similarity). SUBCELLULAR LOCATION: Type I membrane protein. SIMILARITY: Contains 3 immunoglobulin-like C2-type domains. SIMILARITY: Contains 1 immunoglobulin-like V-type domain.
                                                                                                                                                                                                                                                                                                                                                                                                            D86589; BAA13132.1; -..., X73322; CAA51748.1; -..., X703226; AAB60873.1; -...; AF001228; AAB60875.1; -...; AF057380; AAC25124.1; -...; P01730; 1WIQ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SWISS-PROT entry is copyright. It is produced through en the Swiss Institute of Bioinformatics and the EN
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                                                                                                                                                                                                                                  CD4TCANTIGEN.
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                                                                                                                                                                                                       response;
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        I; T-cell; Transmembrane.

By similarity.

By similarity.

T-cell surface glycoprotein CI
Extracellular (potential).

Cytoplasmic (potential).

Ig-like V-type.

Ig-like C2-type 1.

Ig-like C2-type 2.

Ig-like C2-type 3.

N-linked (GlCNAc. . .) (Potent
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01-OCT-1996 (Re]
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T-cell surface (
This SWISS-PROT entry is copyright. It is produced through a collabbetween the Swiss Institute of Bioinformatics and the EMBL outst the European Bioinformatics Institute. There are no restrictions use by non-profit institutions as long as its content is in modified and this statement is not removed. Usage by and for come entities requires a license agreement (See http://www.isb-sib.ch/anor send an email to licenseash-air.
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CONFLICT
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Fomsgaard A., Hirsch V.M., Johnson P.R.;
"Cloning and sequences of primate CD4 molecule
"Cellular receptor for simian immunodeficiency
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Bukaryota, Metazoa, Chordata,
Mammalia, Eutheria, Primates,
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(Rel. 34, Last sequence update)
(Rel. 44, Last annotation update)
ace glycoprotein CD4 (T-cell surface antigen T4/Leu-3)
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xta; Craniata; Vertebrata; Eutele
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Q1-FEB-1995 (Rel. 31, Last secondary)
Q5-JUL-2004 (Rel. 44, Last and
T-cell surface glycoprotein C
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SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
Glycoprotein; Immune response; Immunoglobulin domain; Lipoprotein;
Palmitate; Repeat; T-cell; Transmembrane.
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InterPro; IPR00973; CD4 TCA9.
InterPro; IPR007110; Ig-Tike.
InterPro; IPR003596; Ig_v.
Pfam; PP00047; ig; 2.
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GO:0015026; F:coreceptor activity; ISS.
GO:0042289; F:MHC class II protein binding; ISS.
GO:0006955; P:immune response; ISS.
GO:0045086; P:positive regulation of interleukin
GO:00330217; P:T-cell differentiation; ISS.
GO:0045058; P:T-cell selection; ISS.
GO:0045058; P:T-cell selection; ISS.
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X73327; CAA51753.1;
P01730; IWIQ.
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Cytoplasmic (Potential).
Ig-like V-type.
Ig-like C2-type 1.
Ig-like C2-type 2.
Ig-like C2-type 3.
Ig-like C2-type 3.
Ig-like C1cNAc. . .) (IN-linked (GlCNAc. .) (IN-linked (GlCNA
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Pred. No.
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CD4 (T-cell surface antigen
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HSSP; P01730; IWIO.
GO; GO:0042101; C:T-cell recep)
GO; GO:0015026; F:coreceptor an
GO; GO:0042289; F:MHC class II
GO; GO:0045086; P:jammune respo)
GO; GO:0045086; P:positive respo)
GO; GO:0030217; P:T-cell differ
GO; GO:0030217; P:T-cell differ
GO; GO:0045058; P:T-cell differ
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Primates; Catarrhini; Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pomsgaard A., Hirsch V.M., Johnson P.R.; "Cloning and sequences of primate CD4 molecule cellular receptor for simian immunodeficiency
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cercopithecinae; Erythrocebus.
NCBI_TaxID=9538;
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Bur. J. Immunol. 22:2973-2981(1992).
Eur. J. Immunol. 22:2973-2981(1992).
I. FUNCTION: Accessory protein for MHC class-II antigen/T-cell receptor interaction. May regulate T-cell activation.
receptor interaction. May regulate T-cell activation.
I. SUBUNIT: Associates with p56-lck (By similarity).
I. SUBUNIT: Associates wit
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pfam; PF00047; ig; 2.
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InterPro; IPR000973; CD4 TCA9.
InterPro; IPR007110; Ig-Tike.
InterPro; IPR003596; Ig_v.
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palmitate; Re
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GO:0015026; F:coreceptor activity; ISS.
GO:0042289; F:MHC class II protein binding; ISS.
GO:0045086; P:memune response; ISS.
GO:0045086; P:positive regulation of interleukin-2 biosyn.
GO:0030217; P:T-cell differentiation; ISS.
GO:0030217; P:T-cell selection; ISS.
GO:0045058; P:T-cell selection; ISS.
GO:0045058; P:transmembrane receptor protein tyrosine kin.
GO:0007169; P:transmembrane receptor protein tyrosine kin.
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SM00406; IGV;
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Cytoplasmic (Potential).
Ig-like C2-type 1.
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N-linked (GlcMAC. . .) (B)
Semilarity.
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S-palmitoyl Cysteine (By 1)
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Pred. No. 3.5e-57;
0; Mismatches 13
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EMBL; AF001225; AAB60872.1; -.

HSSP; P01730; INTQ.

GO; GO:0016021; C:integral to membrane; IE

GO; GO:0007155; P:cell adhesion; IEA.

GO; GO:0006955; P:immune response; IEA.

InterPro; IPR008424; CD2.

InterPro; IPR00973; CD4 TCA9.

InterPro; IPR0007110; Ig-Tike.

InterPro; IPR000396; Ig_v.

R Pfam; PF00947; ig; 1.

R Pfam; PF00047; ig; 1.

R PRINTS; PR00692; CD4TCANTIGEN.

R PRNATT; SM00406; IGv; 1.

R PROSITE; PS50835; IG_LIKE; 1.

R PROSITE; PS50835; IG_LIKE; 1.
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Best Local Similarity
Matches 148; Conserv
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009261;
01-JUL-1997
01-JUL-1997
01-OCT-2003
 009262;
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Fomegaard A., Muller-Trutwin M.C., Diop O., Hansen J., N
Corbet S., Barre-Sinoussi F., Allan J.S.;
"Relation between phylogeny of African green monkey CD4
their respective simian immunodeficiency virus genes.";
J. Med. Primatol. 26:120-128(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cercopithecus sabaeus.

Bukaryota; Metazoa; Chordata; C
Mammalia; Butheria; Primates; C
Cercopithecinae; Cercopithecus.
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                                                                                                                       GSSPSVQCRSPRGKNIQGGKTLSVSQLELQDSGTWTCTVLQNQKKVEFKIDIV
                                                                                                                                                                   VVLGKKGDTVELTCNASQNTTTQFHWKNSNQIKILGKQGSFLTKGSSKLRDRIDSRKSLW
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397 1
                                                                                                                                                                                                                                                                                                        74.5%; Score 767; DB 2; llarity 85.5%; Pred. No. 7.6e-56; Conservative 10; Mismatches 15
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                   PRELIMINARY;
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Last sequence update)
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                   PRT;
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Best Local (
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01-JUL-1997
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01-JUL-1997
01-OCT-2003
 MEDLINE=98017879; PubMed=9379478; Fomegaard A., Muller-Trutwin M.C., Diop Corbet S., Barre-Sinoussi F., Allan J.S. "Relation between phylogeny of African g
                                                                                                                  Cercopithecus tantalus.

Eukaryota; Metazoa; Chordata; C
Mammalia; Eutheria; Primates; C
Cercopithecinae; Cercopithecus.
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Bukáryota; Metazoa; Chordata; C
Mammalia; Eutheria; Primates; C
Cercopithecinae; Cercopithecus.
                                                                                                                                                                                       Name=CD4;
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                                                                                                                                                                                                                                                                                          009263
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                                                                  SEQUENCE FROM N.A.
                                                                                                  NCBI_TaxID=60712;
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397 AA;
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(TrEMBLrel.
(TrEMBLrel.
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(TrEMBLrel.
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EMBL; AF001221; ABB60868.1; -.
HSSP; P01730; 1WIQ.
GO; GO:0016021; C:integral to membrane; IE
GO; GO:0007155; P:cell adhesion; IEA.
GO; GO:0006955; P:immune response; IEA.
InterPro; IPR008424; CD2.
InterPro; IPR00973; CD4 TCAg.
InterPro; IPR009710; Ig-Ike.
InterPro; IPR009710; Ig-Ike.
InterPro; IPR003596; Ig_v.
Pfam; PF00047; Ig; 1.
PRINTS; PR00406; IGV; 1.
PRINTS; SM00406; IGV; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            43954 MW; CF7F2F5D82335B0D CRC64;
                                                                                                                                                                                                                                                                                                                       Last sequence update)
Last annotation update)
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Last annotation updat
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Pred. No. 7.6e-56;
                                                                                                                                                                                                  Craniata; Vertebrata; Euteleostomi; Catarrhini; Cercopithecidae;
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                                                 Hansen J., Mathiot
  monkey
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CD4 genes
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RESULT 13
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ID Q95NE
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OC ELMAX
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Best Local :
                                                                   InterPro; IPR008424; CD2.
InterPro; IPR000973; CD4 TCA9.
InterPro; IPR00710; Ig-Tike.
InterPro; IPR003596; Ig-V.
Pfam; PF05790; C2-8et; 2.
Pfam; PF00047; ig; 1.
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InterPro; IPR00873; CD4 TCA9.
InterPro; IPR007110; Ig-Tike.
InterPro; IPR007110; Ig-Tike.
InterPro; IPR007110; Ig-Tike.
Pfam; PP05790; C2-set; Z.
Pfam; PP00047; Ig; I.
PRINTS; PR00692; CD4TCANTIGEN.
SMART; SM00406; IGV; I.
PROSITE; PS50835; IG LIKE; I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Q95NE9;
Q95NE9;
01-DEC-2001
01-DEC-2001
01-OCT-2003
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HSSP; P01730; 1W;
GO; GO:0016021; GO; GO:0007155; GO; GO:0007155; GO; GO:0006955;
                 PRINTS; PR00692; CD4TCI
SMART; SM00406; IGV; 1.
                                                                                                                                                                                                        HSSP; P01730; IWIQ.
GO; GO:0016021; C:integral to membrane;
GO; GO:0007155; P:cell adhesion; IEA.
GO; GO:0006955; P:immune response; IEA.
                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.

MEDLINE=98017879; PubMed=9379478;

Fomsgaard A., Muller-Trutwin M.C., Diop O., Hansen J., Mathiot Corbet S., Barre-Sinoussi F., Allan J.S.;

"Relation between phylogeny of African green monkey CD4 genes a their respective simian immunodeficiency virus genes.";

J. Med. Primatol. 26:120-128(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cercopithecus pygerythrus (Vervet monkey).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cercopithecinae; (NCBI_TaxID=60710;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Name=CD4
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d. Primatol. 26:120-128(1997).
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397 AA;
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(TrEMBLrel. 19, Last sequence update)
(TrEMBLrel. 25, Last annotation update)
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                                     ig; 1.
2; CD4TCANTIGEN.
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IG_LIKE; 1
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Pred. No. 7.6e-56;
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InterPro; IPR000973; CD4 TCAg.
InterPro; IPR000973; IP-Tike.
InterPro; IPR003596; Ig-v.
Pfam; PF005790; C2-set; 2.
Pfam; PF00047; Ig; 1.
PRINTS; PR00692; CD4TCANTIGEN.
SWART; SM00406; IG-y; 1.
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Fonmsgaard A., Muller-Trutwin M.C., Diop O., Hansen J.,
Corbet S., Barre-Sinoussi F., Allan J.S.;
"Relation between phylogeny of African green monkey CD-
their respective simian immunodeficiency virus genes.",
J. Med. Primarol. 26:120-128(1997).
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01-JUL-1997 (TrEMBLrel.
01-JUL-1997 (TrEMBLrel.
01-OCT-2003 (TrEMBLrel.
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GO:0007155; P:cell adhesion; T
GO:0006955; P:immune response;
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148; Conser
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                                                                                    VVLGKKGDTVELTCNASQKTTTQFHWKNSNQIKILGKQGSFLTKGSSKLRDRIDSRKSLW
                                                                                                           VVLGKKGDTVELTCTASQKKSIQFHWKNSNQIKILGNQGSFLTKGFSKLNDRADSRRSLW
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DOGCESMITKNIKIEDSETYICEVENKEBEVELLVEGITANSDTHLLOGOSLTLTLESPP
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397 AA;
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Pred. No. 1.7e-55;
L1; Mismatches 15;
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Pred. No. 7.6e-56;
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RESULT 15
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1D 009260
1D 009260;
1D 009260;
1D 01-JUL-1997 (TREMBLrel. 04, Last sequence update)
DT 01-JUL-1997 (TREMBLrel. 25, Last annotation update)
DT 01-JUL-1997 (TREMBLrel. 25, Last annotation update)
DT 01-OCT-2003 (TREMBLREL. 25, Last annotation update)
DE Cot Cercopithecus sabaeus.
OC Bukaryota; Metazoa; Chordata; Craniata; Verrebrata; But Cot Mammalia; Sutheria; Pubmedes; Catarrhini; Cercopithecid OC C
Search completed: March
Job time: 75.9095 secs
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                                                                                                                                                                                                                                          GSSPSVKCRSPRGKNIQGGRTLSVPQLERQDSGTWTCNVSQDQNTVEFKIDIV 173
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